FIGURE 1

1	CTGCAGTGAC	CAUTGCCCCA	TCATTGCTGG	CTGAGGTGGT	TGGGGTCCAT	CTGGCTATCT
61	GGGCAGCTGT	TCTCTTCTCT	CCTTTCTCTC	CTGTTTCCAG	ACATGCAGTA	TTTCCAGAGA
121	GAAGGGGCCA	CTCTTTGGCA	AAGAACCTGT	CTAACTTGCT	ATCTATGGCA	GGACCTTTGA
181	AGGGTTCACA	GGAAGCAGCA	CAAATTGATA	CTATTCCACC	AAGCCATCAG	CTCCATCTCA
241	TCCATGCCCT	GTCTCTCCTT	TAGGGGTCCC	CTTGCCAACA	GAATCACAGA	GGACCAGCCT
301	GAAAGTGCAG	AGACAGCAGC	TGAGGCACAG	CCAAGAGCTC	TGGCTGTATT	AATGACCTAA
361	GAAGTCACCA	GAAAGTCAGA	AGGATGCATA	GCAGAGGCCC	AGCAATCTCA	GCTAAGTCAA
421	CTCCACCAGC	CTITCTAGTT	GCCCACTGTG	TGTACAGCAC	SCTGGTAGGG	ACCAGAGCCA
481	TGACAGGGAA	TAAGACTAGA	CTATGCCCTT	GAGGAGCTCA	CCTCTGTTCA	GGGAAACAGG
541	CGTGGAAACA	CAATGGTGGT	AAAGAGGAAA	GAGGACAATA	GGATTGCATG	AAGGGGATGG
601					GTTTGGTGAG	
	AAGAGAAGGC					
721	GGCACACTCC	AGGCATAGGT	AAAGATCTGT	AGGTGTGGCT	TGTTGGGATG	AATTTCAAGT
	ATTTTGGAAT					
841	ATGCCAATGG	CTCCACTTGA	GTTTCTGATA	AGAACCCAGA	ACCCTTGGAC	TCCCCAGTAA
	0					
	GTGTGTGTGT					
	AGCACAGGTG					
	ACAATCCAAC					
	CCCAGCAAAG					
	AGTAGTGATG					
1261	CCTGGTGCTC	CTCTATCTGT	GAGTAACTGT	TCAG GCTCCT	CTTCTCTGTT	TCTTGGACTT
1321	GGGGTCGTAA	TCAGGCCTCT	CTTTT			

FIGURE 2

1	GATCTCAGAT	ATCCCTTCTA	TCTACACATT	ATCTATAATT	CTTTCTTTCT	GTAAACTGAA
61	AGGTCCTAGA	AGGAGCCGCA	GCTCAGCAGG	AGAGAGGAGG	AGCTGAGCTG	GGACCCCTAC
121	CTCCTGAGGA	ATGAAATGAT	TATTATAAAG	ACAGCAACCG	AGCTTATTTT	ACCCAAAATA
181	AGGTAGTATA	TTTCTGTTAG	AGTTTAGAGT	TTCATGAGTC	AGGGACCAAG	TTATTGCTTT
241	TCTTTGCCCT	GTATAAAGGC	TTCTCCAAGG	CCTTTGACTT	ACCTAAGTAC	TAAATGTTAT
301	AAAACCAAAC	TCTTCTGACC	TCTCAATCTA	GTCAACTGGG	GCTGTAATTA	TTAATGAAAT
361	TAATGTTTAT	TTTGAAAATA	ATTTACTAGA	CTGAATTACG	AAATCCTGAA	TCATTGTACA
421	CTATCAGTAA	ATATTGGTGG	ACCCAACTGA	ACTGAATGTT	TTGCTTGAAA	TGAAACCTTT
	GAGATGCAGG					
	GCTAAGATAC					
	GTAAGCAGAG					
661	TGTTAGAATC	CCTGTTAAAA	ATGACCAGTA	AAGCTT TGTG	CAACTGTGTC	TTGACATAAC
721	TTTATTTTTC	TTAATAAAAG	AAATGGAAAT	AACCTCACTA	GGGAATTTAG	AACAAATATG
	${\tt ATGATATCTT}$					
	TATCTTTCTA					
	CATTGTGGTG					
	TGCATCCACA					
	TATCTGAAAA					
	GTCGAGAAGC					
	CGTTTCACTT					
1201	GGTGCTGTTT	CCCTTAGAGA	CAAATAAGGG	${\tt GTTCTATTTA}$	ATGTGAAGCC	TGTTTTATGA
	ACAGAATAAA					
1321	TAAAAATTGT	TCTCAAGGCA	GCTCTGGTGT	AAGAGA TAAT	ACACCACGAT	GGGCATCAGA
1381	AGACCTCAGC	TCAAATCCCA	GTTCTGCCAG	CTATGAGCTG	TGTGGCACCA	ACAGGTGTCC
1441	TGTTCTCCCA	GGGTCTCCCT	TTTCCCATTT	GAAAAATAAA	AAATAACAAT	TCCTGCCTTC
1501	AGGAATTTTT	TTTAGGGGGT	TTAATKGTAA	${\tt AGGTGTTTAT}$	ATCTGCTAAG	GTAATTTACT
	TGATATATGT					
	CTGTATTTTT					
	TCCTAATCTT					
	ATTTCTGTTA					
	AATTGCTTTT					
	CATCTGGGTT					
	ACTGGAATGT					
	TTTATCTGTA					
	TGGTCAATTT					
	TGGAGTGCAA					
	GTCCTTGTGC					
	AGAGGAAAAC					
	GGTATTAAGG					
	AAAGGTAAGT				ATCCCTAGAG	GTACATGTTA
2401	CAAGAGGTAA	TGGTAAAGTA	AAATACTTTG	AAAGGCTT		
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